

Could you verify their identity-- or else advise us to destroy them. If the strain is as correctly described in your ms., just check it off; many of them are not specifically cited. (In general, I would advise giving strain numbers in detail in publications, to allow others to make specific citations of their origin and identity). The strains are as follows: (Why not just return this sheet)

J2F+

* Pairing: it would seem that all (B-)M- lines have a fairly high degree of "pairing" even when one is not selecting for this chromosome, while T.B.- lines have low "pairing". ~~all~~ Pairing is variably from experiment to experiment, so that this result is not ~~clearly~~ clear cut.